



L Number	Hits	Search Text	DB	Time stamp
1	13512	coli and (swine or pig or cattle)	USPAT;	2003/09/30 11:06
			US-PGPUB;	
			EPO;	
_			DERWENT	0000/00/00
2	4944	, ,- ,- ,- ,- ,- ,- ,-	USPAT;	2003/09/30 11:06
		(intestine or intestinal)	US-PGPUB;	
			EPO;	
			DERWENT	2002/00/20 11:07
3	57		USPAT;	2003/09/30 11:07
		(intestine or intestinal) and (FUT? or	US-PGPUB; EPO;	
		fucosyltransferase)	DERWENT	
4	2	(coli and (swine or pig or cattle) and	USPAT;	2003/09/30 11:07
.   *	2	(intestine or intestinal) and (FUT? or	US-PGPUB;	2003/03/30 11:07
		fucosyltransferase)).clm.	EPO;	
			DERWENT	
5	10	coli and (swine or pig or cattle) and	USPAT;	2003/09/30 11:08
-		(intestine or intestinal) and (FUT? or	US-PGPUB;	
		fucosyltransferase) and (alteration or	EPO;	
-		mutation or polymorphism).clm.	DERWENT	

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Leu Val Cys Val Leu Ala Ala Ile Phe Phe Leu Asn Val Tyr Gln Asp
                                          25
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ctc ttt tac agt ggc tta gac ctg ctg gcc ctg tgt cca gac cat aac
                                                                    146
Leu Phe Tyr Ser Gly Leu Asp Leu Leu Ala Leu Cys Pro Asp His Asn
                 35
gtg gta tca tct ccc gtg gcc ata ttc tgc ctg gcg ggc acg ccg gta
Val Val Ser Ser Pro Val Ala Ile Phe Cys Leu Ala Gly Thr Pro Val
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cac His	ccc	aac Asn 65	Ala	tcc Ser	gat Asp	tcc Ser	tgt Cys _70	Pro	aag Lys	g cat His	cct Pro	gcc Ala 75	Ser	ttt Phe	tcc Ser	242
ggg ggg	acc Thr 80	Trp	act Thr	att Ile	tac Tyr	ecg Pro 85	gat Asp	ggc	cgg	ttt Phe	999 Gly	Asn	cag Gln	atc Met	gga Gly	290
cag Gln 95	tat Tyr	gcc Ala	acg Thr	ctg Leu	ctg Leu 100	gcc Ala	ctg Leu	gcg Ala	cag Gln	ctc Leu 105	Asn	ggc	cgc Arg	cag Gln	gcc Ala 110	338
ttc Phe	atc Ile	cag Gln	cct Pro	gcc Ala 115	Met	cac His	gcc Ala	gtc Val	ctg Leu 120	gcc Ala	ccc Pro	gtg Val	ttc Phe	cgc Arg 125	Ile	386
acg Thr	ctg Leu	cct Pro	gtc Val 130	ctg Leu	gcg Ala	ccc Pro	gag Glu	gta Val 135	gac Asp	agg Arg	cac His	gct Ala	cct Pro 140	tgg Trp	cgg Arg	434
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cac His 175	cac His	ctc Leu	cgg Arg	gag Glu	cag Gln 180	atc Ile	cgc Arg	agc Ser	gag Glu	ttc Phe 185	acc Thr	ctg Leu	cac His	gac Asp	cac His 190	578
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ggc Gly	cgt Arg 240	tac Tyr	ctc Leu	cag Gln	cag Gln	gct Ala 245	atg Met	gac Asp	tgg Trp	ttc Phe	cgg Arg 250	gcc Ala	cga Arg	tac Tyr	gaa Glu	770
gcc Ala 255	ccc Pro	gtc Val	ttt Phe	Val	gtc Val 260	acc Thr	agc Ser	aac Asn	Gly	atg Met 265	gag Glu	tgg Trp	tgc Cys	cgg Arg	aag Lys 270	818
aac Asn	atc Ile	gac Asp	acc Thr	tcc Ser 275	cgg Arg	ggg Gly	gac Asp	Val	atc Ile 280	ttt Phe	gct Ala	ggc Gly	gat Asp	999 Gly 285	cgg Arg	866

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Glu Ala Ala Pro Ala Arg Asp Phe Ala Leu Leu Val Gln Cys Asn His 290 . 295 300	. 914
acc atc atg acc att ggc acc ttc ggc ttc tgg gcc gcc tac ctg gct Thr Ile Met Thr Ile Gly Thr Phe Gly Phe Trp Ala Ala Tyr Leu Ala 305 310 315	962
ggt gga gat acc atc tac ttg gct aac ttc acc ctg ccc act tcc agc Gly Gly Asp Thr Ile Tyr Leu Ala Asn Phe Thr Leu Pro Thr Ser Ser 320 325 330	1010
ttc ctg aag atc ttt aaa ccc gag gct gcc ttc ctg ccc gag tgg gtg Phe Leu Lys Ile Phe Lys Pro Glu Ala Ala Phe Leu Pro Glu Trp Val 335 340 345 350	1058
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360